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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2011; month=3; day=8; hr=6; min=43; sec=28; ms=690;]

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Application No: 10552291 Version No: 2.0

Input Set:

Output Set:

Started: 2011-03-07 16:54:26.216
Finished: 2011-03-07 16:54:32.175
Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 959 ms
Total Warnings: 47
Total Errors: 0
No. of SeqIDs Defined: 57
Actual SeqID Count: 57

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 213	Artificial or Unknown found in <213> in SEQ ID (23)
W 213	Artificial or Unknown found in <213> in SEQ ID (24)
W 213	Artificial or Unknown found in <213> in SEQ ID (25)
W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)
W 213	Artificial or Unknown found in <213> in SEQ ID (28)
W 213	Artificial or Unknown found in <213> in SEQ ID (29)
W 213	Artificial or Unknown found in <213> in SEQ ID (30)

Input Set:

Output Set:

Started: 2011-03-07 16:54:26.216
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Actual SeqID Count: 57

Error code	Error Description
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SEQUENCE LISTING

<110> REGEN Biotech, Inc.

<120> Use of a peptide that interacts with alpha v beta3 integrin of endothelial cell

<130> OP04-1024

<140> 10552291

<141> 2005-10-03

<150> KR 10-2003-0021065

<151> 2003-04-03

<160> 57

<170> KopatentIn 1.71

<210> 1

<211> 683

<212> PRT

<213> Homo sapiens

<400> 1

Met Ala Leu Phe Val Arg Leu Leu Ala Leu Ala Leu Ala Leu
1 5 10 15

Gly Pro Ala Ala Thr Leu Ala Gly Pro Ala Lys Ser Pro Tyr Gln Leu
20 25 30

Val Leu Gln His Ser Arg Leu Arg Gly Arg Gln His Gly Pro Asn Val
35 40 45

Cys Ala Val Gln Lys Val Ile Gly Thr Asn Arg Lys Tyr Phe Thr Asn
50 55 60

Cys Lys Gln Trp Tyr Gln Arg Lys Ile Cys Gly Lys Ser Thr Val Ile
65 70 75 80

Ser Tyr Glu Cys Cys Pro Gly Tyr Glu Lys Val Pro Gly Glu Lys Gly
85 90 95

Cys Pro Ala Ala Leu Pro Leu Ser Asn Leu Tyr Glu Thr Leu Gly Val
100 105 110

Val Gly Ser Thr Thr Gln Leu Tyr Thr Asp Arg Thr Glu Lys Leu
115 120 125

Arg Pro Glu Met Glu Gly Pro Gly Ser Phe Thr Ile Phe Ala Pro Ser
130 135 140

Asn Glu Ala Trp Ala Ser Leu Pro Ala Glu Val Leu Asp Ser Leu Val
145 150 155 160

Ser Asn Val Asn Ile Glu Leu Leu Asn Ala Leu Arg Tyr His Met Val
165 170 175

Gly Arg Arg Val Leu Thr Asp Glu Leu Lys His Gly Met Thr Leu Thr
180 185 190

Ser Met Tyr Gln Asn Ser Asn Ile Gln Ile His His Tyr Pro Asn Gly
195 200 205

Ile Val Thr Val Asn Cys Ala Arg Leu Leu Lys Ala Asp His His Ala
210 215 220

Thr Asn Gly Val Val His Leu Ile Asp Lys Val Ile Ser Thr Ile Thr
225 230 235 240

Asn Asn Ile Gln Gln Ile Ile Glu Ile Glu Asp Thr Phe Glu Thr Leu
245 250 255

Arg Ala Ala Val Ala Ala Ser Gly Leu Asn Thr Met Leu Glu Gly Asn
260 265 270

Gly Gln Tyr Thr Leu Leu Ala Pro Thr Asn Glu Ala Phe Glu Lys Ile
275 280 285

Pro Ser Glu Thr Leu Asn Arg Ile Leu Gly Asp Pro Glu Ala Leu Arg
290 295 300

Asp Leu Leu Asn Asn His Ile Leu Lys Ser Ala Met Cys Ala Glu Ala
305 310 315 320

Ile Val Ala Gly Leu Ser Val Glu Thr Leu Glu Gly Thr Thr Leu Glu
325 330 335

Val Gly Cys Ser Gly Asp Met Leu Thr Ile Asn Gly Lys Ala Ile Ile
340 345 350

Ser Asn Lys Asp Ile Leu Ala Thr Asn Gly Val Ile His Tyr Ile Asp
355 360 365

Glu Leu Leu Ile Pro Asp Ser Ala Lys Thr Leu Phe Glu Leu Ala Ala
370 375 380

Glu Ser Asp Val Ser Thr Ala Ile Asp Leu Phe Arg Gln Ala Gly Leu
385 390 395 400

Gly Asn His Leu Ser Gly Ser Glu Arg Leu Thr Leu Leu Ala Pro Leu
405 410 415

Asn Ser Val Phe Lys Asp Gly Thr Pro Pro Ile Asp Ala His Thr Arg
420 425 430

Asn Leu Leu Arg Asn His Ile Ile Lys Asp Gln Leu Ala Ser Lys Tyr
435 440 445

Leu Tyr His Gly Gln Thr Leu Glu Thr Leu Gly Gly Lys Lys Leu Arg
450 455 460

Val Phe Val Tyr Arg Asn Ser Leu Cys Ile Glu Asn Ser Cys Ile Ala
465 470 475 480

Ala His Asp Lys Arg Gly Arg Tyr Gly Thr Leu Phe Thr Met Asp Arg
485 490 495

Val Leu Thr Pro Pro Met Gly Thr Val Met Asp Val Leu Lys Gly Asp
500 505 510

Asn Arg Phe Ser Met Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr
515 520 525

Glu Thr Leu Asn Arg Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn
530 535 540

Glu Ala Phe Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly
545 550 555 560

Asp Ala Lys Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu
565 570 575

Ile Leu Val Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu
580 585 590

Gln Gly Asp Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val
595 600 605

Asn Lys Glu Pro Val Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val
610 615 620

Val His Val Ile Thr Asn Val Leu Gln Pro Pro Ala Asn Arg Pro Gln
625 630 635 640

Glu Arg Gly Asp Glu Leu Ala Asp Ser Ala Leu Glu Ile Phe Lys Gln
645 650 655

Ala Ser Ala Phe Ser Arg Ala Ser Gln Arg Ser Val Arg Leu Ala Pro
660 665 670

Val Tyr Gln Lys Leu Leu Glu Arg Met Lys His
675 680

<210> 2

<211> 103

<212> PRT

<213> Homo sapiens

<400> 2

Gly Pro Gly Ser Phe Thr Ile Phe Ala Pro Ser Asn Glu Ala Trp Ala
1 5 10 15

Ser Leu Pro Ala Glu Val Leu Asp Ser Leu Val Ser Asn Val Asn Ile
20 25 30

Glu Leu Leu Asn Ala Leu Arg Tyr His Met Val Gly Arg Arg Val Leu
35 40 45

Thr Asp Glu Leu Lys His Gly Met Thr Leu Thr Ser Met Tyr Gln Asn

50	55	60
Ser Asn Ile Gln Ile His His Tyr Pro Asn Gly Ile Val Thr Val Asn		
65	70	75
Cys Ala Arg Leu Leu Lys Ala Asp His His Ala Thr Asn Gly Val Val		
85	90	95
His Leu Ile Asp Lys Val Ile		
100		

<210> 3
<211> 131
<212> PRT
<213> Homo sapiens

<400> 3		
Asn Ile Gln Gln Ile Ile Glu Ile Glu Asp Thr Phe Glu Thr Leu Arg		
1	5	10
15		

Ala Ala Val Ala Ala Ser Gly Leu Asn Thr Met Leu Glu Gly Asn Gly		
20	25	30

Gln Tyr Thr Leu Leu Ala Pro Thr Asn Glu Ala Phe Glu Lys Ile Pro		
35	40	45

Ser Glu Thr Leu Asn Arg Ile Leu Gly Asp Pro Glu Ala Leu Arg Asp		
50	55	60

Leu Leu Asn Asn His Ile Leu Lys Ser Ala Met Cys Ala Glu Ala Ile		
65	70	75
80		

Val Ala Gly Leu Ser Val Glu Thr Leu Glu Gly Thr Thr Leu Glu Val		
85	90	95

Gly Cys Ser Gly Asp Met Leu Thr Ile Asn Gly Lys Ala Ile Ile Ser		
100	105	110

Asn Lys Asp Ile Leu Ala Thr Asn Gly Val Ile His Tyr Ile Asp Glu		
115	120	125

Leu Leu Ile
130

<210> 4
<211> 129
<212> PRT
<213> Homo sapiens

<400> 4		
Pro Asp Ser Ala Lys Thr Leu Phe Glu Leu Ala Ala Glu Ser Asp Val		
1	5	10
15		

Ser Thr Ala Ile Asp Leu Phe Arg Gln Ala Gly Leu Gly Asn His Leu		
20	25	30

Ser Gly Ser Glu Arg Leu Thr Leu Leu Ala Pro Leu Asn Ser Val Phe
35 40 45

Lys Asp Gly Thr Pro Pro Ile Asp Ala His Thr Arg Asn Leu Leu Arg
50 55 60

Asn His Ile Ile Lys Asp Gln Leu Ala Ser Lys Tyr Leu Tyr His Gly
65 70 75 80

Gln Thr Leu Glu Thr Leu Gly Lys Lys Leu Arg Val Phe Val Tyr
85 90 95

Arg Asn Ser Leu Cys Ile Glu Asn Ser Cys Ile Ala Ala His Asp Lys
100 105 110

Arg Gly Arg Tyr Gly Thr Leu Phe Thr Met Asp Arg Val Leu Thr Pro
115 120 125

Pro

<210> 5

<211> 131

<212> PRT

<213> Homo sapiens

<400> 5

Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
1 5 10 15

Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
20 25 30

Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
35 40 45

Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
50 55 60

Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val Ser Gly
65 70 75 80

Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp Lys Leu
85 90 95

Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu Pro Val
100 105 110

Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val Val His Val Ile Thr
115 120 125

Asn Val Leu
130

<210> 6
<211> 85
<212> PRT
<213> Homo sapiens

<400> 6
Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys
1 5 10 15

Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val
20 25 30

Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp
35 40 45

Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu
50 55 60

Pro Val Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val Val His Val
65 70 75 80

Ile Thr Asn Val Leu
85

<210> 7
<211> 119
<212> PRT
<213> Homo sapiens

<400> 7
Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
1 5 10 15

Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
20 25 30

Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
35 40 45

Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
50 55 60

Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val Ser Gly
65 70 75 80

Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp Lys Leu
85 90 95

Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu Pro Val
100 105 110

Ala Glu Pro Asp Ile Met Ala
115

<210> 8

<211> 113
<212> PRT
<213> Homo sapiens

<400> 8
Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
1 5 10 15

Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
20 25 30

Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
35 40 45

Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
50 55 60

Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val Ser Gly
65 70 75 80

Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp Lys Leu
85 90 95

Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu Pro Val
100 105 110

Ala

<210> 9
<211> 73
<212> PRT
<213> Homo sapiens

<400> 9
Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys
1 5 10 15

Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val
20 25 30

Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp
35 40 45

Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu
50 55 60

Pro Val Ala Glu Pro Asp Ile Met Ala
65 70

<210> 10
<211> 67
<212> PRT
<213> Homo sapiens

<400> 10
Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys
1 5 10 15

Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val
20 25 30

Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp
35 40 45

Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu
50 55 60

Pro Val Ala
65

<210> 11
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-AA(18)

<400> 11
Lys Glu Leu Ala Asn Ile Leu Lys Ala Ala Ile Gly Asp Glu Ile Leu
1 5 10 15

Val Ser

<210> 12
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-L(18)

<400> 12
Lys Glu Ser Ala Asn Ser Ser Lys Tyr His Ile Gly Asp Glu Ile Leu
1 5 10 15

Val Ser

<210> 13
<211> 18
<212> PRT
<213> Artificial Sequence

<220>

<223> D-IV-R(18)

<400> 13
Lys Glu Leu Ala Asn Ile Leu Lys Tyr His Ser Gly Asp Glu Ser Ser
1 5 10 15

Val Ser

<210> 14
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-LYHR(18)

<400> 14
Lys Glu Ser Ala Asn Ser Ser Lys Tyr His Ser Gly Asp Glu Ser Ser
1 5 10 15

Val Ser

<210> 15
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-LAA(18)

<400> 15
Lys Glu Ser Ala Asn Ser Ser Lys Ala Ala Ile Gly Asp Glu Ile Leu
1 5 10 15

Val Ser

<210> 16
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-AAR(18)

<400> 16
Lys Glu Leu Ala Asn Ile Leu Lys Ala Ala Ser Gly Asp Glu Ser Ser
1 5 10 15

Val Ser

<210> 17
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-AA

<400> 17
Gly Asp Ala Lys Glu Leu Ala Asn Ile Leu Lys Ala Ala Ile Gly Asp
1 5 10 15

Glu Ile Leu Val Ser Gly Gly Ile Gly Ala Leu Val Arg
20 25

<210> 18
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-L

<400> 18
Gly Asp Ala Lys Glu Ser Ala Asn Ser Ser Lys Tyr His Ile Gly Asp
1 5 10 15

Glu Ile Leu Val Ser Gly Gly Ile Gly Ala Leu Val Arg
20 25

<210> 19
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-R

<400> 19
Gly Asp Ala Lys Glu Leu Ala Asn Ile Leu Lys Tyr His Ser Gly Asp
1 5 10 15

Glu Ser Ser Val Ser Gly Gly Ile Gly Ala Leu Val Arg
20 25

<210> 20
<211> 29

<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-LYHR

<400> 20
Gly Asp Ala Lys Glu Ser Ala Asn Ser Ser Lys Tyr His Ser Gly Asp
1 5 10 15

Glu Ser Ser Val Ser Gly Gly Ile Gly Ala Leu Val Arg
20 25

<210> 21
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-LAA

<400> 21
Gly Asp Ala Lys Glu Ser Ala Asn Ser Ser Lys Ala Ala Ile Gly Asp
1 5 10 15

Glu Ile Leu Val Ser Gly Gly Ile Gly Ala Leu Val Arg
20 25

<210> 22
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> D-IV-AAR

<400> 22
Gly Asp Ala Lys Glu Leu Ala Asn Ile Leu Lys Ala Ala S